

SEQUENCE LISTING

<110> Reiner, Peter B.
 Connop, Bruce P.
 Pollard, Michelle

<120> REGULATION OF AMYLOID PRECURSOR PROTEIN EXPRESSION
 BY MODIFICATION OF ABC TRANSPORTER EXPRESSION OR ACTIVITY

<130> 100103.402

<140> US

<141> 2002-02-08

<160> 10

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 3512

<212> DNA

<213> Homo sapiens

<400> 1

cgccccgggca	ggtcagcctg	tctcaaggca	cgccagtcctc	agctccgacc	ttgcagcggc	60
gcagcgcggg	tgggagggcg	ggaggagcag	cggaagagc	ggagcgagga	cccggtccgg	120
cgcagtcctc	aatgagcagc	gcggaactg	caccccagac	ccgagcctgc	tgcgcgcccc	180
ctcccagagc	tcacctggtg	ccaggttaaca	ggcctggcct	cgccctgtgg	atgatgatgg	240
ccttgccccc	gtgagctaca	acctggcctt	cagcaccgc	ccacctccaa	ccagcaggat	300
gcggctgtgg	aaggcggtgg	tggtgacttt	ggccttcctg	agtgtggaca	tctgcgtgac	360
cacggccatc	tatgtcttca	gccacctgga	ccgcagcctc	ctggaggaca	tccgccactt	420
caacatcttt	gactcggtgc	tggatctctg	ggcagcctgc	ctgtaccgca	gctgcctgct	480
gctgggagcc	accattggtg	tggccaagaa	cagtgcgctg	gggccccggc	ggctgcgggc	540
ctcgtggctg	gtcatcacc	tcgtgtgcct	cttcgtgggc	atctatgcca	tgggtgaagct	600
gctgctcttc	tcagaggtgc	gcaggcccat	ccgggacccc	tgggttttggg	ccctgttcgt	660
gtggacgtac	atttcaactg	gcgcctcctt	cctgctctgg	tggctgctgt	ccaccgtgcg	720
gccaggcacc	caggccctgg	agccaggggc	ggccaccgag	gctgagggct	tccctgggag	780
cggccggcca	ccgcccgagc	aggcgtctgg	ggccacgctg	cagaagctgc	tctcctacac	840
caagcccgc	gtggccttcc	tcgtggccgc	ctccttcttc	ctcatcgtgg	cagctctggg	900
agagaccttc	ctgccctact	acacgggccc	cgccattgat	ggcatcgtca	tccagaaaag	960
catggatcag	ttcagcacgg	ctgtcgtcat	cgtgtgcctg	ctggccattg	gcagctcatt	1020
tgcgcaggt	attcggggcg	gcatttttac	cctcatattt	gccagactga	acattcgcct	1080
tcgaaactgt	ctcttcgcgt	cactgggtgc	caggagaca	agcttctttg	atgagaaccg	1140
cacaggggac	ctcatctccc	gcctgacctc	ggacaccacc	atggtcagcg	acctggtctc	1200
ccagaacatc	aatgtcttcc	tgcggaacac	agtcaaggctc	acgggcgctg	tggctctcat	1260
gttcagcctc	tcattggcagc	tctccttggg	caccttcctg	ggcttcccca	tcattcatgat	1320
ggtgtccaac	atctacggca	agtactacaa	gaggctctcc	aaagaggctc	agaatgccct	1380
ggccagagcg	agcaacacgg	cggaggagac	catcagtgcc	atgaagactg	tccggagctt	1440
cgccaatgag	gaggaggagg	cagaggtgta	cctgcggaag	ctgcagcagg	tgtacaagct	1500
gaacaggaag	gaggcagctg	cctacatgta	ctacgtctgg	ggcagcgggc	tcacactgct	1560
ggtggtccag	gtcagcatcc	tctactacgg	gggccacctt	gtcatctcag	gccagatgac	1620

cagcggcaac	ctcatcgcct	tcatcatcta	cgagtttgtc	ctgggagatt	gtatggagtc	1680
cgtgggctcc	gtctacagt	gcctgatgca	gggagtggg	gctgctgaga	aggtgttcga	1740
gttcatcgac	cggcagccga	ccatgggtga	cgatggcagc	ttggccccc	accacctgga	1800
gggcccgggtg	gactttgaga	atgtgacctt	cacctaccgc	actcgccccc	acacccaggt	1860
cctgcagaat	gtctccttca	gcctgtcccc	cggcaagggtg	acggccctgg	tggggccctc	1920
gggcagtggg	aagagctcct	gtgtcaacat	cctggagaac	ttctaccccc	tggagggggg	1980
ccgggtgctg	ctggacggca	agcccatcag	cgctacgac	cacaagtact	tgcaccgtgt	2040
gatctccctg	gtgagccagg	agcccggtgt	gttcgcccgc	tccatcacgg	ataacatctc	2100
ctacggcctg	cccactgtgc	ctttcgagat	ggtggtggag	gccgcacaga	aggccaatgc	2160
ccacggcttc	atcatggaac	tccaggacgg	ctacagcaca	gagacagggg	agaagggcgc	2220
ccagctgtca	ggtggccaga	agcagcgggt	ggccatggcc	cgggctctgg	tgcggaaccc	2280
cccagtcctc	atcctggatg	aagccaccag	cgctttggat	gccgagagcg	agtatctgat	2340
ccagcaggcc	atccatggca	acctgcagaa	gcacacggta	ctcatcatcg	cgcaaccggt	2400
gagcaccgtg	gagcacgcgc	acctcattgt	ggtgctggac	aagggccgcg	tagtgacgca	2460
gggcacccac	cagcagctgc	tggcccaggg	cggcctctac	gccaagctgg	tgcagcggca	2520
gatgctgggg	cttcagcccc	ccgcagactt	cacagctggc	cacaacgagc	ctgtagccaa	2580
cggcagtcac	aaggcctgat	ggggggcccc	tgtttctccc	ggtggggcag	aggaccgggt	2640
gcctgcctgg	cagatgtgcc	cacggaggcc	cccagctgcc	ctccgagccc	aggcctgcag	2700
cactgaaaga	cgacctgcca	tgtcccatgg	atcaccgctt	cctgcatctt	gcccctgggtc	2760
cctgccccat	tcccagggca	ctccttacc	ctgctgccct	gagccaaacgc	cttcacggac	2820
ctccctagcc	tcctaagcaa	aggtagagct	gcctttttta	acctaggtct	taccagggtt	2880
tttactgttt	ggtttgaggg	accccagtc	actcctagat	ttcaaaaacc	tttttctaat	2940
tgggagtaat	ggcgggcact	ttcaccaaga	tgttctagaa	acttctgagc	caggagtga	3000
tggcccttcc	ttagtagcct	gggggatgtc	cagagactag	gcctctcccc	tttaccctc	3060
cagagaaggg	gcttccctgt	cccggaggga	cacggggaac	gggattttcc	gtctctccct	3120
cttgccagct	ctgtgagtct	ggccaggggc	ggtaggagc	gtggagggca	tctgtctgcc	3180
atcgcccgct	gccaatctaa	gccagtctca	ctgtgaacca	cacgaaacct	caactggggg	3240
agtgaagggc	tggccaggtc	tggaggggcc	tcagggggtg	ccagcccggc	accagcgct	3300
ttcgccctc	gtccacccac	cccgtgctgg	cagcctccct	cccacacccc	gcccctgtgc	3360
tctgtgtct	ggaggccacg	tggatgttca	tgagatgcat	tctcttctgt	ctttgggtgga	3420
tgggatggtg	gcaaagccca	ggatctggct	ttgccagagg	ttgcaacatg	ttgagagaac	3480
ccggtcaata	aagtgtacta	cctcttacc	ct			3512

<210> 2

<211> 4643

<212> DNA

<213> Homo sapiens

<400> 2

cctactctat	tcagatatto	tccagattcc	taaagattag	agatcatttc	tcattctcct	60
aggagtactc	acttcaggaa	gcaaccagat	aaaagagagg	tgcaacggaa	gccagaacat	120
tcctcctgga	aattcaacct	gtttcgcagt	ttctcgagga	atcagcattc	agtcaatccg	180
ggccgggagc	agtcatctgt	ggtgaggctg	attggctggg	caggaacagc	gccggggcgt	240
gggctgagca	cagcgcttcg	ctctcttttc	cacaggaagc	ctgagctcat	tcgagtagcg	300
gctcttccaa	gtctcaagaa	gcagaggccg	ctgttcgttt	cctttagggtc	tttccactaa	360
agtcggagta	tcttcttcca	agatttcacg	tcttggtggc	cgttccaagg	agcgcgaggt	420
cgggatggat	cttgaagggg	accgcaatgg	aggagcaaag	aagaagaact	tttttaaact	480
gaacaataaa	agtgaaaaag	ataagaagga	aaagaaacca	actgtcagtg	tattttcaat	540
gtttcgctat	tcaaattggc	ttgacaagtt	gtatatggtg	gtgggaactt	tggctgccat	600
catccatggg	gctggacttc	ctctcatgat	gctgggtgtt	ggagaaatga	cagatatctt	660
tgcaaataga	ggaaatttag	aagatctgat	gtcaaacatc	actaatagaa	gtgatataca	720
tgatacaggg	ttcttcatga	atctggagga	agacatgacc	aggtatgcct	attattacag	780
tggaaattgg	gctgggggtg	tgggtgctgc	ttacattcag	gtttcatttt	ggtgcctggc	840
agctggaaga	caaatacaca	aaattagaaa	acagtttttt	catgctataa	tgcgacagga	900

gataaggctgg	tttgatgtgc	acgatgttgg	ggagcttaac	acccgactta	cagatgatgt	960
ctccaagatt	aatgaaggaa	ttggtgacaa	aattggaatg	ttctttcagt	caatggcaac	1020
attttttact	gggtttatag	taggattttac	acgtggttgg	aagctaacc	ttgtgatttt	1080
ggccatcagt	cctgttcttg	gactgtcagc	tgctgtctgg	gcaaagatac	tatcttcatt	1140
tactgataaa	gaactccttag	cgtatgcaaa	agctggagca	gtagctgaag	aggtccttggc	1200
agcaattaga	actgtgattg	catttgagg	acaaaagaaa	gaacttgaaa	ggtacaacaa	1260
aaatttagaa	gaagctaaaa	gaattgggat	aaagaaagct	attacagcca	atatttctat	1320
aggtgctgct	ttcctgctga	tctatgcata	ttatgctctg	gccttctggt	atgggaccac	1380
cttggctctc	tcaggggaa	attctattgg	acaagtactc	actgtatttt	ctgtattaat	1440
tggggctttt	agtgttggac	aggcatctcc	aagcattgaa	gcatttgcaa	atgcaagagg	1500
agcagcttat	gaaatcttca	agataattga	taataagcca	agtattgaca	gctattcgaa	1560
gagtgggcac	aaaccagata	atattaaggg	aaatttggaa	ttcagaaatg	ttcacttcag	1620
ttacccatct	cgaaaagaag	ttaagatctt	gaagggtctg	aacctgaagg	tgcagagtgg	1680
gcagacggtg	gccctggttg	gaaacagtgg	ctgtgggaag	agcacaacag	tccagctgat	1740
gcagaggctc	tatgacccca	cagaggggat	ggtcagtggt	gatggacagg	atattaggac	1800
cataaattga	aggtttctac	gggaaatcat	tggtgtggtg	agtcaggaa	ctgtattggt	1860
tgccaccacg	atagctgaaa	acattcgcta	tggcctgtaa	aatgtcacca	tggatgagat	1920
tgagaaagct	gtcaaggaag	ccaatgccta	tgactttatc	atgaaactgc	ctcataaatt	1980
tgacaccctg	gttgagagag	gaggggcccc	gttgagtgg	gggcagaagc	agaggatcgc	2040
cattgcacgt	gccctggttc	gcaaccccaa	gatcctctctg	ctggatgagg	ccacgtcagc	2100
cttgacaca	gaaagcgaag	cagtggttca	ggtggctctg	gataaggcca	gaaaaggctg	2160
gaccaccatt	gtgatagctc	atcgtttgtc	tacagttcgt	aatgctgacg	tcacgtctgg	2220
tttcgatgat	ggagtcat	tggagaaagg	aatcatgat	gaactcatga	aagagaaagg	2280
catttacttc	aaacttgtca	caatgcagac	agcaggaaat	gaagtgaat	tagaaaatgc	2340
agctgatgaa	tccaaaagt	aaattgatgc	cttggaatg	tcttcaaattg	attcaagatc	2400
cagtctaata	agaaaaagat	caactcgtag	gagtgtccgt	ggatcacaa	ccaagacag	2460
aaagcttagt	accaaagagg	ctctggatga	aagtatacct	ccagtttct	tttggaggat	2520
tatgaagcta	aatttaactg	aatggcctta	ttttgttgtt	ggtgtatttt	gtgccattat	2580
aatggaggc	ctgcaaccag	catttgcaat	aatattttca	aagattatag	gggtttttac	2640
aagaattgat	gatcctgaaa	caaaacgaca	gaatagtaac	ttgttttcac	tattgtttct	2700
agcccttga	attatttctt	ttattacatt	tttcttcag	ggtttcacat	ttggcaaagc	2760
tggagagatc	ctcaccaagc	ggctccgata	catggttttc	cgatccatgc	tcagacagga	2820
tgtgagttgg	tttcatgacc	ctaaaaacac	cactggagca	ttgactacca	ggctcgccaa	2880
tgtatctgct	caagttaaag	gggcttatag	ttccaggctt	gctgtaatta	cccagaatat	2940
agcaaatctt	gggcagagaa	taattatata	cttctctat	ggttgccaac	taacactggt	3000
actcttagca	attgtaccca	tcattgcaat	agcaggagtt	gttgaaatga	aaatgttgtc	3060
tggacaagca	ctgaaagata	agaaagaact	agaagggtct	gggaagatcg	ctactgaagc	3120
aatagaaaa	ttccgaaccg	ttgtttcttt	gactcaggag	cagaagtttg	aacatatgta	3180
tgctcagagt	ttgcaggtag	catacagaaa	ctctttgagg	aaagcacaca	tctttggaat	3240
tacattttcc	ttcaccacag	caatgatgta	tttttctat	gctggatgtt	tcoggtttgg	3300
agcctacttg	gtggcacata	aactcatgag	ctttgaggat	gttctgttag	tattttcagc	3360
tgttgtcttt	ggtgccatgg	ccgtggggca	agtcagttca	tttgtctctg	actatgccaa	3420
agccaaaata	tcagcagccc	acatcatcat	gatcattgaa	aaaaccctt	tgattgacag	3480
ctacagcacg	gaaggcctaa	tgccgaacac	attggaagga	aatgtcacat	ttggtgaagt	3540
tgtattcaac	tatcccaccc	gaccggacat	cccagtgcct	cagggaactga	gcctggagg	3600
gaagaagggc	cagacgctgg	ctctggtggg	cagcagtggc	tgtgggaaga	gcacagtggt	3660
ccagctcctg	gagcggttct	acgacccctt	ggcaggga	gtgctgcttg	atggcaaaga	3720
aataaagcga	ctgaatgttc	agtggctccg	agcacacctg	ggcatcgtgt	cccaggagcc	3780
catcctgttt	gactgcagca	ttgctgagaa	cattgcctat	ggagacaaca	gccgggtgg	3840
gtcacaggaa	gagattgtga	gggcagcaaa	ggaggccaac	atacatgcct	tcacagagtc	3900
actgccta	aaatatgca	ctaaagttag	agacaaagga	actcagctct	ctggtggcca	3960
gaaacaacgc	attgccatag	ctcgtgccct	tggttagacg	ctcatatttt	tgttttggga	4020
tgaagccacg	tcagctctgg	atacagaaa	tgaaaaggtt	gtccaagaag	ccctggacaa	4080
agccagagaa	ggccgcacct	gcatttgtat	tgctcaccgc	ctgtccacca	tccaqaatgc	4140

agacttaata	gtggtggttc	agaatggcag	agtcaaggag	catggcacgc	atcagcagct	4200
gctggcacag	aaaggcatct	atTTTTcaat	ggtcagtgtc	caggctggaa	caaagcgcca	4260
gtgaactctg	actgtatgag	atgttaaata	ctttttaata	tttgtttaga	tatgacattt	4320
attcaaagtt	aaaagcaaac	acttacagaa	ttatgaagag	gtatctgttt	aacatttcct	4380
cagtcaagtt	cagagtcttc	agagacttcg	taattaaagg	aacagagtga	gagacatcat	4440
caagtggaga	gaaatcatag	tttaaactgc	attataaatt	ttataacaga	attaaagtag	4500
atTTTaaaag	ataaaatgtg	taattttgtt	tatatTTTcc	catttggact	gtaactgact	4560
gccttgctaa	aagattatag	aagtagcaaa	aagtattgaa	atgtttgcat	aaagtgtcta	4620
taataaaaact	aaactttcat	gtg				4643

<210> 3

<211> 8056

<212> DNA

<213> Homo sapiens

<400> 3

cgcgagaggg	agcgggcgcg	ggcgctgagg	cgcgaggagcg	tggccccgcc	atgggcttcc	60
tgcaccagct	gcagctgctg	ctctggaaga	acgtgacgct	caaacgcggg	agcccggtggg	120
tcttgccctt	cgagatcttc	atccccctgg	tgtgttctt	tatcctgctg	gggctgcgac	180
agaagaagcc	cacctctctc	gtgaaggaag	tccccctcta	cacagcgggc	cccctgacgt	240
ctgccggcat	cctgcctgtc	atgcaatcgc	tgtgcccgga	cgccagcgga	gacgagttcg	300
gcttcctgca	gtacgccaac	tccacggtca	cgcagctgct	tgagcgctg	gaccgctggg	360
tggaggaagg	caacctgttt	gaccagcgcg	ggccagcct	gggtcagag	ctcgaggccc	420
tacgccagca	tctggaggcc	ctcagtgcgg	gcccgggcac	ctcggggagc	cacctggaca	480
gatccacagt	gtcttctctc	tctctggact	cggtggccag	aaaccgcgag	gagctctggc	540
gtttcctgac	gcaaaaacttg	tcgctgcccc	atagcacggc	ccaagcactc	ttggccgccc	600
gtgtggaccc	gcccagagtc	taccactcgc	tctttggtcc	ctcatctgcc	ctggattcac	660
agtctggcct	ccacaagggt	caggagccct	ggagccgcct	agggggcaat	cccctgttcc	720
ggatggagga	gctgctgctg	gctcctgccc	tcttgagca	gctcacctgc	acgcggggt	780
cgggggagct	gggcccggatc	ctcactgtgc	ctgagagtca	gaaggagagc	ctgcagggt	840
accgggatgc	tgtctgcagt	gggcaggctg	ctgcgcgtgc	caggcgcttc	tctgggctgt	900
ctgctgagct	ccggaaccag	ctggacgtgg	ccaaggtctc	ccagcagctg	ggcctggatg	960
cccccaacgg	ctcggaactc	tcgccacagg	cgccaccccc	acggaggctg	caggcgcttc	1020
tgggggacct	gctgggatgc	cagaagggtc	tgcaggatgt	ggatgtcctg	tcggccctgg	1080
ccctgctact	gccccagggt	gcctgcaactg	gccggacccc	cggacccccca	gccagtgggtg	1140
cggttggggc	ggccaatggc	actggggcag	gggcagtcac	gggcccccaac	gccaccgctg	1200
aggaggggcg	accctctgct	gcagcactgg	ccaccccgga	cacgctgcag	ggccagtgt	1260
cagccttcgt	acagctctgg	gccggcctgc	agcccatctt	gtgtggcaac	aaccgcacca	1320
ttgaaccoga	ggcgtgctgg	cggggcaaca	tgagctccct	gggttccacg	agcaaggagc	1380
agcggaacct	gggcctcctc	gtgcacctca	tgaccagcaa	ccccaaaatc	ctgtacgcgc	1440
ctgcgggctc	tgaggctcgac	cgctcatc	tcaaggccaa	cgagactttt	gcttttgtgg	1500
gcaacgtgac	tcactatgcc	caggtctggc	tcaacatctc	ggcggagatc	cgcagcttcc	1560
tggagcaggg	caggctgcag	caacacctgc	gctggctgca	gcagtatgta	gcagagctgc	1620
ggctgcaccc	cgaggcactg	aacctgtcac	tggatgagct	gccgcgggcc	ctgagacagg	1680
acaacttctc	gctgccaggt	ggcatggccc	tcttgacgca	gctggatacc	attgacaacg	1740
cggcctgcgg	ctggatccag	ttcatgtcca	agtgagcgt	ggacatcttc	aagggttcc	1800
ccgacgagga	gagcattgtc	aactacaccc	tcaaccaggc	ctaccaggac	aacgtcactg	1860
tttttgccag	tgtgatcttc	cagaccggga	aggacggctc	gctcccgct	cacgtgcact	1920
acaagatccg	ccagaactcc	agcttcacgg	agaaaaccaa	cgagatccgc	cgcgcctact	1980
ggcggcctgg	gcccataact	ggcggcgcgt	tctacttcct	ctacggcttc	gtctggatcc	2040
aggacatgat	ggagcgcgcc	atcatcgaca	cttttgtggg	gcacgacgtg	gtggagccag	2100
gcagctacgt	gcagatgttc	ccctaccct	gctacacacg	cgatgacttc	ctgtttgtca	2160
ttgagcacat	gatgccgctg	tgcattggtg	tctcctgggt	ctactcctg	gccatgacca	2220
tccagcacat	cgtggcggag	aaggagcacc	ggctcaaggga	ggtgatgaag	accatggggc	2280

tgaacaacgc	ggtgcaactgg	gtggcctggt	tcatcacccg	ctttgtgcag	ctgtccatct	2340
ccgtgacagc	actcacgcgc	atcctgaagt	acggccaggt	gcttatgcac	agccacgtgg	2400
tcatcatctg	gctcttccctg	gcagtctacg	cgggtggccac	catcatgttc	tgcttccctgg	2460
tgtctgtgct	gtactccaag	gccaaactgg	cctcggcctg	cgggtggcatc	atctacttcc	2520
tgagctacgt	gccctacatg	tacgtggcga	tccgagagga	ggtggcgcac	gataagatca	2580
cggccttcga	gaagtgcac	gcgtccctca	tgccacgcac	ggcctttggt	ctgggctcta	2640
agtacttcgc	gctgtatgag	gtggccggcg	tgggcatcca	gtggcacacc	ttcagccagt	2700
ccccgggtga	gggggacgac	ttcaacttgc	tccctggctgt	caccatgctg	atggtggacg	2760
ccgtggtcta	tggcatcctc	acgtggtaca	ttgaggctgt	gcaccaggc	atgtacgggc	2820
tgccccggcc	ctggtacttc	ccactgcaga	agtccactg	gctgggcagt	gggcggacag	2880
aagcctggga	gtggagctgg	ccgtgggcac	gcaccccccg	cctcagtgct	atggaggagg	2940
accaggcctg	tgccatggag	agccggcgct	ttgaggagac	ccgtggcatg	gaggaggagc	3000
ccacccacct	gcctctggtt	gtctgcgtgg	acaaactcac	caaggtctac	aaggacgaca	3060
agaagctggc	cctgaacaag	ctgagcctga	acctctacga	gaaccagggtg	gtctccttct	3120
tgggccacaa	cggggcgggc	aagaccacca	ccatgtccat	cctgaccggc	ctgttccctc	3180
caacgtcggg	ttccgccacc	atctacgggc	acgacatccg	cacggagatg	gatgagatcc	3240
gcaagaacct	gggcatgtgc	ccgcagcaca	atgtgctctt	tgaccggctc	acggtggagg	3300
aacacctctg	gttctactca	cggctcaaga	gcatggctca	ggaggagatc	cgcagagaga	3360
tggacaagat	gatcgaggac	ctggagctct	ccaacaaacg	gcactcactg	gtgcagacat	3420
tgtcgggtgg	catgaagcgc	aagctgtccg	tggccatcgc	cttcgtgggc	ggctctcgcg	3480
ccatcatcct	ggacgagccc	acggcgggcg	tggaccctta	cgcgcgcgcg	gccatctggg	3540
acctcatcct	gaagtacaag	ccaggccgca	ccatccttct	gtccacccac	cacatggatg	3600
aggctgacct	gcttggggac	cgcattgcca	tcatctccca	tgggaagctc	aagtgtctgcg	3660
gtcccccgct	cttctcaag	ggcacctatg	gcgacgggta	ccgcctcacg	ctggtcaagc	3720
ggcccgccga	gcccggggggc	ccccaaagac	cagggtctggc	atccagcccc	ccaggctcggg	3780
ccccgctgag	cagctgctcc	gagctccagg	tgtcccagtt	catccgcaag	catgtggcct	3840
cctgcctgct	ggtctcagac	acaagcacgg	agctctccta	catcctgccc	agcgaggccg	3900
ccaagaaggg	ggctttcgag	cgcctcttcc	agcacctgga	gcgcagcctg	gatgcactgc	3960
acctcagcag	cttcgggctg	atggacacga	ccctggaggga	agtgttctct	aagggtctcg	4020
aggaggatca	gtcgtctggg	aacagtgagg	ccgatgtgaa	ggagtccagg	aaggatgtgc	4080
tccttggggc	ggagggcccg	gcgtctgggg	agggtcacgc	tggcaatctg	gcccgggtgct	4140
cggagctgac	ccagtgcgag	gcacgtctgc	agtcggcgct	atctgtgggc	tctgcccgtg	4200
gcgacgaggg	agctggctac	accgacgtct	atggcgacta	ccgccccctc	tttgataacc	4260
cacaggaccc	agacaatgtc	agcctgcaag	agggtggaggc	agaggccctg	tcgagggtcg	4320
gccagggcag	ccgcaagctg	gacggcgggg	ggctgaagggt	gcgccagttc	cacgggctgc	4380
tgggtcaaacg	cttccactgc	gcccgcgcga	actccaaggc	actcttctcc	cagatcttgc	4440
tgcacgcctt	cttcgtctgc	gtggccatga	ccgtggccct	gtccgtcccg	gagattggtg	4500
atctgcccc	gctggtcctg	tcaccttccc	agtaccacaa	ctacaccag	ccccgtggca	4560
atttcatccc	ctacgccaac	gaggagcgcc	gcgagtaacc	gctgcggcta	tcgcccagcg	4620
ccagccccc	gcagctcgtg	agcacgttcc	ggctgccgtc	gggggtgggt	gccacctgcg	4680
tgtcaagtc	tcccgccaac	ggctcgctgg	ggccccagtt	gaacctgagc	agcggggagt	4740
cgcgcctgct	ggcggtcggg	ttcttcgaca	gcatgtgtct	ggagtcttct	acacaggggc	4800
tgccactgtc	caatttcgtg	ccacccccac	cctcgcccg	cccatctgac	tcgccagcgt	4860
ccccggatga	ggacctgcag	gcctggaacg	tctccctgcc	gcccaccgct	gggccagaaa	4920
tgtggacgtc	ggcacctctc	ctgcgcgcgc	tggtaacggga	gcccgtccgc	tgcacctgct	4980
ctgcgcaggg	caccggcttc	tctgccccca	gcagtgtggg	cgggcacccg	cccagatgc	5040
gggtggtcac	aggcgacatc	ctgaccgaca	tcaccggcca	caatgtctct	gagtacctgc	5100
tcttcacctc	cgaccgcttc	cgactgcacc	ggatatggggc	catcaccttt	ggaaacgtcc	5160
tgaagtccat	cccagcctca	tttggcacca	gggccccacc	catggtgcgg	aagatcgcg	5220
tgcgcagggc	tgcccaggtt	ttctacaaca	acaagggtta	tcacagcatg	cccacctacc	5280
tcaacagcct	caacaacgcc	atcctgcgtg	ccaacctgcc	caagagcaag	ggcaacccgg	5340
cggcttacgg	catcacgctc	accaaccacc	ccatgaataa	gaccagcgcc	agcctctccc	5400
tggattacct	gctgcagggc	acggatgtcg	tcatcgccat	cttcatcatc	gtggccatgt	5460
ccttcgtgcc	ggccagcttc	gttgtcttcc	tctgtggcca	gaagtccacc	aaggccaagc	5520

```

atctgcagtt tgtcagcggc tgcaacccca tcatctactg gctggcgaac tacgtgtggg 5580
acatgctcaa ctacctggto cccgctacct gctgtgtcat catcctgttt gtgttcgacc 5640
tgccggccta cagctgcgcc accaacttcc ctgccgtcct ctccctcttc ctgctctatg 5700
ggtggtccat cagcoccate atgtaccceg cctccttctg gttcgaggte cccagctccg 5760
cctacgtgtt cctcattgtc atcaatctct tcatcgcat caccgccacc gtggccacct 5820
tcctgctaca gctcttcgag cagacaagg acctgaagg tgtcaacagt tacctgaaaa 5880
gctgcttcc cttttcccc aactacaacc tgggccacgg gctcatggag atggcctaca 5940
acgagtacat caacgagtac tacgccaaga ttggccagtt tgacaagatg aagtcctcgt 6000
tcgagtggga cattgtcacc cgcggactgg tggccatggc ggttgagggc gtcgtgggct 6060
tcctcctgac catcatgtgc cagtacaact tcctgcgggc gccacagcgc atgcctgtgt 6120
ctaccaagcc tgtggaggat gatgtggacg tggccagtga gcggcagcga gtgctccggg 6180
gagacgccga caatgacatg gtcaagattg agaacctgac caaggtctac aagtcctcga 6240
agattggccg tatectggcc gttgaccgcc tgtgcctggg tgtgcgtcct ggagagtgt 6300
tcgggctcct gggcgtcaac ggtgcgggca agaccagcac cttcaagatg ctgaccggcg 6360
acgagagcac gacggggggc gaggccttgc tcaatggaca cagcgtgctg aaggagctgc 6420
tcaggtgca gcagagcctc ggctactgcc cgcagtgtga cgcgtgttc gacgagctca 6480
cggcccgga gcacctgcag ctgtacacgc ggctgcgtgg gatctcctgg aaggacgagg 6540
cccgggtggt gaagtgggct ctggagaagc tggagctgac caagtacgca gacaagccgg 6600
ctggcaccta cagcggcggc aacaagcgga agctctccac ggccatcgcc ctattgggt 6660
accagcctt catcttctctg gacgagccca ccacaggcat ggaccccaag gcccggcgt 6720
tcctctgga cctcatctc gacctcatca agacaggcg ttcagtgggt ctgacatcac 6780
acagcatgga ggagtgcgag gcgctgtgca cgcgctggc catcatgggt aacggtcgcc 6840
tgcggtgctt gggcagcatc cagcacctga agaaccggtt tggagatggc tacatgatca 6900
cggtcgggac caagagcagc cagagtgtga aggacgtgtt gcggttcttc aaccgcaact 6960
tcccggaagc catgctcaag gagcggcacc acacaaagg tgcagtaccag ctcaagtcgg 7020
agcacatctc gctggcccag gtgttcagca agatggagca ggtgtctggc gtgctgggca 7080
tcgaggacta ctcggtcagc cagaccacac tggacaatgt gttcgtgaac tttgccaaga 7140
agcagagtga caacctggag cagcaggaga cggagccgcc atccgactg cagtccctc 7200
tcggtgctt gctcagcctg ctccggcccc ggtctgcccc caccgagctc cgggcacttg 7260
tggcagacga gcccaggagc ctggacacgg aggacgaggg cctcatcagc ttcgaggagg 7320
agcgggcccc gctgtccttc aacacggaca cgtctgctg accaaccaga gctgggccag 7380
ggaggacacg ctccactgac caccagagc tgggccaggg actcaacaat ggggacagaa 7440
gtccccagc gctgccagg gcctggagt gaggttcagg accaaggggc ttctggtcct 7500
ccagccctg tactcgcca tgccctgcgg tcatgcggt tgcgcctcct aattgtgcca 7560
aaggctgacc cggcccgggc tgcgtacacc cttgcctgc tttgccttaa agcctcgggg 7620
tctgccggc cctcgcccc tgctggcac tgctcaccgc ccaaggcgac gccggtgga 7680
ccaggcactg ctggcctttc tcctgcccgg cctcggaacc agcttttctc tcttacgatg 7740
aaggctgatg ccgagagcgg gctgtgggag gagctgggtc agtcccgtat ttattttgct 7800
ttgagaagag gctcctctgg cctgctctc ctgcaggag gtggctgtcc cgcgggaagc 7860
catcagcttg ggccagctgg caggtggcag gaatggagaa gctgacctg ctggccaggc 7920
aaggggcccag acccccccca acccccagct gccatcgctc tcccaccag cttggcccc 7980
tgcccgcaca cctccttggg agccgggct gtacatagcg cacagatgtt tgttttaaat 8040
aaataaacia aatgtc 8056

```

<210> 4

<211> 3455

<212> DNA

<213> Homo sapiens

<400> 4

```

gccacatgg cggagaaggc gctggaggcc gtgggctgtg gactagggcc gggggctgtg 60
gccatggcgg tgacgttga ggacggggcg gaacccctg tgctgaccac gcacctgaag 120
aaggtggaga accacatcac tgaagcccag cgcttctccc acctgccaa gcgctcagcc 180

```

gtggacatcg	agttcgtgga	gctgtcctat	tccgtgcggg	aggggccctg	ctggcgcaaa	240
aggggttata	agacccttct	caagtgcctc	tcaggtaa	tctgccgcg	ggagctgatt	300
ggcatcatgg	gccccctcagg	ggctggcaag	tctacattca	tgaacatctt	ggcaggatac	360
agggagtctg	gaatgaaggg	gcagatcctg	gttaatggaa	ggccacggga	gctgaggacc	420
ttccgcaaga	tgtcctgcta	catcatgcaa	gatgacatgc	tgtcgccgca	cctcaagggtg	480
ttggaagcca	tgatgggtctc	tgctaacctg	aatcttactg	agaatcccga	tgtgaaaaac	540
gatctcgtga	cagagatcct	gacggcactg	ggcctgatgt	cgtgctccca	cacgaggaca	600
gccctgctct	ctggcgggca	gaggaagcgt	ctggccatcg	ccctggagct	ggtcaacaac	660
ccgctgtca	tgttctttga	tgagcccacc	agtggctctg	atagcgctc	ttgtttccaa	720
gtggtgtccc	tcatgaagtc	cctggcacag	gggggccgta	ccatcatctg	caccatccac	780
cagcccagtg	ccaagctctt	tgagatgttt	gacaagctct	acatcctgag	ccagggtcag	840
tgcattctca	aaggagtgg	caccaacctg	atccccatc	taaagggact	cggcttgcat	900
tgccccacct	accacaaccc	ggctgacttc	atcatcgagg	tggcctctgg	cgagtatgga	960
gacctgaacc	ccatgtttgt	cagggctgtg	cagaatgggc	tgtgcgctat	ggctgagaag	1020
aagagcagcc	ctgagaagaa	cgaggctcct	gccccatgcc	ctccttgctc	tcgggaagtg	1080
gatcccatgg	aaagccacac	ctttgccacc	agcaccctca	cacagttctg	catcctcttc	1140
aagaggacct	tcctgtccat	cctcagggac	acggctcctga	cccacctacg	gttcattgtc	1200
cacgtgggta	ttggcggtg	catcggcctc	ctctacctgc	atattggcga	cgatgccagc	1260
aaggctcttca	acaacacccg	ctgcctcttc	ttctccatgc	tgttcctcat	gttogccgcc	1320
ctcatgccaa	ctgtgctcac	cttccccctta	gagatggcgg	tcttcatgag	ggagcacctc	1380
aactactggt	acagcctcaa	agcgtattac	ctggccaaga	ccatggtctga	cgtgcccttt	1440
caggtgggtg	gtccgggtgg	ctactgcagc	atttgttact	ggatgacggg	ccagcccgtc	1500
gagaccagcc	gcttccctgt	cttctcagcc	ctggccaccg	ccaccgcctt	ggtggcccaa	1560
tctttggggc	tgtgatccg	agctgcttcc	aactccctac	aggtggccac	ttttgtgggc	1620
ccagttaccg	ccatccctgt	cctctgttct	tcgggttct	ttgtcagctt	caagaccatc	1680
cccacttacc	tgcaatggag	ctcctatctc	tccatgtca	ggtatggctt	tgagggtgtg	1740
atcctgacga	tctatggcat	ggagcgagga	gacctgacat	gtttagagga	acgtgcccgc	1800
ttccggggagc	cacagagcat	cctccgagcg	ctggatgtgg	aggatgccaa	gctctacatg	1860
gaacttctgg	tcttgggcat	cttcttctca	gcctgcggc	tgttgcccta	ccttgtgctg	1920
cgttaccggg	tcaagtcaga	gagatagagg	cttgccccag	cctgtacccc	agcccctgca	1980
gcaggaagcc	cccagtccca	gccctttggg	actgttttaa	ccttatagac	ttgggcaactg	2040
gttccctggcg	gggctatcct	ctcctccctt	ggctcctcca	caggctggct	gtcggactgc	2100
gctcccagcc	tgggctctgg	gagtgggggc	tccagccctc	cccactatgc	ccaggagtct	2160
tcccaagttg	atgcggtttg	tagcttcctc	cctactctct	ccaacacctg	catgcaaaga	2220
ctactgggag	gctgctgcct	ccttcctgcc	catggcacc	tcctctgctg	tctgcctggg	2280
agccctaggg	tctctagggc	cccacttaca	actgaccaaa	gtggccccct	ctgggggtcc	2340
ccaccacaca	agtgtttgta	aactgggctg	ctataagggt	ggagtccag	ggctggggcc	2400
tggtggagtc	cactggaagt	cccattatgg	atgttgaaat	ggacagggaa	ggactctgga	2460
agtctcttcc	tcctcctcct	cttctctcca	cccctagacc	ctggtgact	tggaacatct	2520
gccaggacag	aagctgggtt	ttctgtctag	gtcaccactc	ccaatcctgg	ggattggaga	2580
ggcctggggc	tgtgggatgc	cccatcccc	tccccatcac	ctttggtggg	ggcagggcct	2640
ggtggcaact	gtgcaataat	gtctgtgttt	ctctcccacc	tgccactgga	actggagaat	2700
gcactttatt	ctgggcgggg	ggtgagtggg	ggaagaccac	accctccttt	ctcgtgccc	2760
ctaacgcattg	cacggctctg	tgatgctccc	tccctctccg	gagtgcagag	cacatacatg	2820
agaacaggcc	atctcagccc	tacacacttg	ccatccccta	cagcacagag	gaagagtgat	2880
ggtggcatgc	tgggtggggc	gggtgctggt	gggaggacag	tgccaacctc	ctcctgggga	2940
tcccatgttg	gagactctaa	ggataaggct	ggtgctgcc	aggggtgtcta	caggaaactgc	3000
aggtgtctac	ccccaaagtct	tcctcctctc	caagccaggg	gtggcacagg	gcactagatc	3060
cctggagtctc	aggaaccaac	acaagcacia	ccacgggcat	aagttggcct	tgggcactgc	3120
cacccacggc	cctccttttg	tgtccatgc	tggcatcttc	actcccctac	cccttcccca	3180
gccactgctg	ctcattcaaa	cttctgtcca	tgtccctcca	ctgttctctat	cagcagggtgg	3240
ccctggggca	tcagaacagc	ctgccctggg	caccagggtgg	cagacacact	cagagcatgt	3300
ctggctttcc	tgggtgggtcc	aggctcattc	tgtctctgat	tccccctccc	ccagggtcca	3360
ttttccccct	ttttcctgta	cacatccctg	tctacctcct	ctcaccctgc	cacagattct	3420

tcctatcaca cagggatgcc agttgtatatt gtggg

3455

<210> 5

<211> 3201

<212> DNA

<213> Homo sapiens

<400> 5

```

gaattccggg atgtggaacg gtcgcaggag gctgctacaa gccccatgag caaggctgtt 60
ccactgaca gagctttccc aggatgacag agagtgcgct ctgcctctct ggggtgtgct 120
agcctacgag gggcaatcgt aaggcgaatg tcaactgaaag aacacaagtg tccttaaaca 180
tggaactatct gggcttttcta gtgctgaaat tcttcccact cccactgccc acttcccatt 240
atataaaaaa cacagttgtt tcatgttttt gtttctttac tgtttttctt tgtttttgtt 300
aagaatgcat tcattttatc aaaattgttt attgtagaat aatcaggcat tgcgtggatg 360
aggtgggtgtc cagcaacatg gaggccactg agacggacct gctgaatgga catctgaaaa 420
aagtagataa taacctcacg gaagcccagc gcttctcctc cttgcctcgg agggcagctg 480
tgaacattga attcagggac ctttcctatt cggttcctga aggaccctgg tggaggaaga 540
aaggatacaa gaccctcctg aaaggaatct ccgggaagtt caatagtggg gagttgggtg 600
ccattatggg tccttcctgg gccgggaagt ccacgctgat gaacatcctg gctggataca 660
gggagacggg catgaagggg gccgtcctca tcaacggcct gccccgggac ctgcgctgct 720
tcgggaaggt gtctctgtac atcatgcagg atgacatgct gctgccgat ctactgtgc 780
aggaggccat gatgggtgtc gcacatctga agcttcagga gaaggatgaa ggcagaaggg 840
aaatgggtcaa ggagatactg acagcgctgg gcttgctgtc ttgcgccaac acgcggaacc 900
ggagcctgtc aggtgggtcag cgcaagcgcc tggccatcgc gctggagctg gtgaacaacc 960
ctccagtcac gttcttcgat gagcccacca gcggcctgga cagcgctcc tgcttccagg 1020
tggtctcgct gatgaaaggg ctcgctcaag ggggtcgctc catcatttgc accatccacc 1080
agcccagcgc caaactcttc gagctgttcg accagcttta cgtcctgagt caaggacaat 1140
gtgtgtaccg gggaaaagtc tgcaatcttg tgccatattt gagggatttg ggtctgaact 1200
gccaaccta ccacaacca gcagattttg tcatggaggt tgcatccggc gactacgggt 1260
atcagaacag tcggctggtg agagcggttc gggaggggcat gtgtgactca gaccacaaga 1320
gagacctcgg ggggtgatgc gaggtgaacc cttttctttg gcaccggccc tctgaagagg 1380
taaagcagac aaaacgatta aaggggttga gaaaggactc ctgctccatg gaaggctgcc 1440
acagcttctc tgccagctgc ctacgcagc tctgcatcct cttcaagagg accttctca 1500
gcatcatgag ggactcggtc ctgacacacc tgcgcatcac ctgcacatt gggatcggcc 1560
tcctcatttg cctgctgtac ttggggatcg ggaacgaagc caagaaggtc ttgagcaact 1620
ccggttctct cttcttctcc atgctgttcc tcatgttcgc ggccctcatg cctactgttc 1680
tgacatttcc cctggagatg ggagtcttcc ttccgggaaca cctgaactac tggtagagcc 1740
tgaaggccta ctacctggcc aagaccatgg cagacgtgcc ctttcagatc atgttcccag 1800
tggcctactg cagcatcgtg tactggatga cgtcgcagcc gtccgacgcc gtggcctttg 1860
tgctgtttgc cgcgctgggc accatgacct cctggtggc acagtccctg ggctgctga 1920
tcggagccgc ctccacgtcc ctgcaggtgg ccactttcgt gggcccagtg acagccatcc 1980
cggtgctcct gttctcgggg ttcttcgtca gcttcgacac catccccacg tacctacagt 2040
ggatgtccta catctcctat gtcaggtatg ggttcgaagg ggtcattctc tccatctatg 2100
gcttagaccg ggaagatctg cactgtgaca tcgacgagac gtgccacttc cagaagtcgg 2160
aggccatcct gcgggagctg gacgtggaag atgccaaagt gtacctggac ttcacgtgac 2220
tcgggatttt cttcatctcc ctccgcctca ttgcctattt tgcctcagg taaaaaatcc 2280
gggcagagag gtaaaacacc tgaatgccag gaaacaggaa gattagacac tgtggccgag 2340
ggcacgtcta gaatcgagga ggcaagcctg tgcccagacc acgacacaga gactcttctg 2400
atccaacccc tagaaccgcg ttgggtttgt ggggtgtctc tgctcagcca ctctgccag 2460
ctgggttgga tcttctctcc attccccttt ctagctttta ctaggaagat gtaggcagat 2520
tggtggtttt ttttttttta acatacagaa ttttaaatac cacaactggg gcagaattta 2580
aagctgcaac acagctggtg atgagaggct tcctcagtc agtcgctcct tagcaccagg 2640
caccgtgggt cctggatggg gaactgcaag cagcctctca gctgatgctg cgcagtcaga 2700

```



```

tgtctggtgg cagagagtcg gagcatggag cgattccatt ttatgactgt tgtttttcaac 2760
attttcatct ttctaagggtg tgtctctttt ccaatgagaa gtcatttttg caagccaaaa 2820
gtcgatcaat cgcattcatt ttaagaaatt atacctttt agtacttget gaagaatgat 2880
tcagggtaaa tcacatactt tgtttagaga ggcgaggggt ttaaccgagt caccagctg 2940
gtctcataca tagacagcac ttgtgaagga ttgaatgcag gttccagggtg gaggaagac 3000
gtggacacca tctccactga gccatgcaga cttttttaa agctatacaa aaaattgtga 3060
gaagacattg gccaaactct tcaaagtctt tctttttcca cgtgcttctt attttaagcg 3120
aaatatattg tttgtttctt cctaaaaacg gaattctttt gctttttacc ctggaagaaa 3180
tactcataat agtagtagta g 3201

```

<210> 6

<211> 766

<212> PRT

<213> Homo sapiens

<400> 6

```

Met Arg Leu Trp Lys Ala Val Val Val Thr Leu Ala Phe Met Ser Val
 1          5          10          15
Asp Ile Cys Val Thr Thr Ala Ile Tyr Val Phe Ser His Leu Asp Arg
 20          25          30
Ser Leu Leu Glu Asp Ile Arg His Phe Asn Ile Phe Asp Ser Val Leu
 35          40          45
Asp Leu Trp Ala Ala Cys Leu Tyr Arg Ser Cys Leu Leu Gly Ala
 50          55          60
Thr Ile Gly Val Ala Lys Asn Ser Ala Leu Gly Pro Arg Arg Leu Arg
 65          70          75          80
Ala Ser Trp Leu Val Ile Thr Leu Val Cys Leu Phe Val Gly Ile Tyr
 85          90          95
Ala Met Val Lys Leu Leu Leu Phe Ser Glu Val Arg Arg Pro Ile Arg
 100         105         110
Asp Pro Trp Phe Trp Ala Leu Phe Val Trp Thr Tyr Ile Ser Leu Gly
 115         120         125
Ala Ser Phe Leu Leu Trp Trp Leu Leu Ser Thr Val Arg Pro Gly Thr
 130         135         140
Gln Ala Leu Glu Pro Gly Ala Ala Thr Glu Ala Glu Gly Phe Pro Gly
 145         150         155         160
Ser Gly Arg Pro Pro Pro Glu Gln Ala Ser Gly Ala Thr Leu Gln Lys
 165         170         175
Leu Leu Ser Tyr Thr Lys Pro Asp Val Ala Phe Leu Val Ala Ala Ser
 180         185         190
Phe Phe Leu Ile Val Ala Ala Leu Gly Glu Thr Phe Leu Pro Tyr Tyr
 195         200         205
Thr Gly Arg Ala Ile Asp Gly Ile Val Ile Gln Lys Ser Met Asp Gln
 210         215         220
Phe Ser Thr Ala Val Val Ile Val Cys Leu Leu Ala Ile Gly Ser Ser
 225         230         235         240
Phe Ala Ala Gly Ile Arg Gly Gly Ile Phe Thr Leu Ile Phe Ala Arg
 245         250         255
Leu Asn Ile Arg Leu Arg Asn Cys Leu Phe Arg Ser Leu Val Ser Gln
 260         265         270
Glu Thr Ser Phe Phe Asp Glu Asn Arg Thr Gly Asp Leu Ile Ser Arg
 275         280         285
Leu Thr Ser Asp Thr Thr Met Val Ser Asp Leu Val Ser Gln Asn Ile
 290         295         300

```

Asn	Val	Phe	Leu	Arg	Asn	Thr	Val	Lys	Val	Thr	Gly	Val	Val	Val	Phe
305					310					315					320
Met	Phe	Ser	Leu	Ser	Trp	Gln	Leu	Ser	Leu	Val	Thr	Phe	Met	Gly	Phe
				325					330						335
Pro	Ile	Ile	Met	Met	Val	Ser	Asn	Ile	Tyr	Gly	Lys	Tyr	Tyr	Lys	Arg
			340					345						350	
Leu	Ser	Lys	Glu	Val	Gln	Asn	Ala	Leu	Ala	Arg	Ala	Ser	Asn	Thr	Ala
		355					360					365			
Glu	Glu	Thr	Ile	Ser	Ala	Met	Lys	Thr	Val	Arg	Ser	Phe	Ala	Asn	Glu
	370					375					380				
Glu	Glu	Glu	Ala	Glu	Val	Tyr	Leu	Arg	Lys	Leu	Gln	Gln	Val	Tyr	Lys
385					390					395					400
Leu	Asn	Arg	Lys	Glu	Ala	Ala	Ala	Tyr	Met	Tyr	Tyr	Val	Trp	Gly	Ser
				405					410						415
Gly	Leu	Thr	Leu	Leu	Val	Val	Gln	Val	Ser	Ile	Leu	Tyr	Tyr	Gly	Gly
			420					425						430	
His	Leu	Val	Ile	Ser	Gly	Gln	Met	Thr	Ser	Gly	Asn	Leu	Ile	Ala	Phe
		435					440					445			
Ile	Ile	Tyr	Glu	Phe	Val	Leu	Gly	Asp	Cys	Met	Glu	Ser	Val	Gly	Ser
	450					455					460				
Val	Tyr	Ser	Gly	Leu	Met	Gln	Gly	Val	Gly	Ala	Ala	Glu	Lys	Val	Phe
465					470					475					480
Glu	Phe	Ile	Asp	Arg	Gln	Pro	Thr	Met	Val	His	Asp	Gly	Ser	Leu	Ala
				485					490						495
Pro	Asp	His	Leu	Glu	Gly	Arg	Val	Asp	Phe	Glu	Asn	Val	Thr	Phe	Thr
			500					505						510	
Tyr	Arg	Thr	Arg	Pro	His	Thr	Gln	Val	Leu	Gln	Asn	Val	Ser	Phe	Ser
		515					520								
Leu	Ser	Pro	Gly	Lys	Val	Thr	Ala	Leu	Val	Gly	Pro	Ser	Gly	Ser	Gly
	530					535					540				
Lys	Ser	Ser	Cys	Val	Asn	Ile	Leu	Glu	Asn	Phe	Tyr	Pro	Leu	Glu	Gly
545					550					555					560
Gly	Arg	Val	Leu	Leu	Asp	Gly	Lys	Pro	Ile	Ser	Ala	Tyr	Asp	His	Lys
				565					570						575
Tyr	Leu	His	Arg	Val	Ile	Ser	Leu	Val	Ser	Gln	Glu	Pro	Val	Leu	Phe
			580					585						590	
Ala	Arg	Ser	Ile	Thr	Asp	Asn	Ile	Ser	Tyr	Gly	Leu	Pro	Thr	Val	Pro
		595					600								605
Phe	Glu	Met	Val	Val	Glu	Ala	Ala	Gln	Lys	Ala	Asn	Ala	His	Gly	Phe
	610					615					620				
Ile	Met	Glu	Leu	Gln	Asp	Gly	Tyr	Ser	Thr	Glu	Thr	Gly	Glu	Lys	Gly
625					630						635				640
Ala	Gln	Leu	Ser	Gly	Gly	Gln	Lys	Gln	Arg	Val	Ala	Met	Ala	Arg	Ala
				645					650						655
Leu	Val	Arg	Asn	Pro	Pro	Val	Leu	Ile	Leu	Asp	Glu	Ala	Thr	Ser	Ala
			660					665						670	
Leu	Asp	Ala	Glu	Ser	Glu	Tyr	Leu	Ile	Gln	Gln	Ala	Ile	His	Gly	Asn
		675					680							685	
Leu	Gln	Lys	His	Thr	Val	Leu	Ile	Ile	Ala	His	Arg	Leu	Ser	Thr	Val
	690					695									
Glu	His	Ala	His	Leu	Ile	Val	Val	Leu	Asp	Lys	Gly	Arg	Val	Val	Gln
705					710					715					720
Gln	Gly	Thr	His	Gln	Gln	Leu	Leu	Ala	Gln	Gly	Gly	Leu	Tyr	Ala	Lys
				725					730						735

Leu Val Gln Arg Gln Met Leu Gly Leu Gln Pro Ala Ala Asp Phe Thr
 740 745 750
 Ala Gly His Asn Glu Pro Val Ala Asn Gly Ser His Lys Ala
 755 760 765

<210> 7
 <211> 1280
 <212> PRT
 <213> Homo sapiens

<400> 7

Met Asp Leu Glu Gly Asp Arg Asn Gly Gly Ala Lys Lys Lys Asn Phe
 1 5 10 15
 Phe Lys Leu Asn Asn Lys Ser Glu Lys Asp Lys Lys Glu Lys Lys Pro
 20 25 30
 Thr Val Ser Val Phe Ser Met Phe Arg Tyr Ser Asn Trp Leu Asp Lys
 35 40 45
 Leu Tyr Met Val Val Gly Thr Leu Ala Ala Ile Ile His Gly Ala Gly
 50 55 60
 Leu Pro Leu Met Met Leu Val Phe Gly Glu Met Thr Asp Ile Phe Ala
 65 70 75 80
 Asn Ala Gly Asn Leu Glu Asp Leu Met Ser Asn Ile Thr Asn Arg Ser
 85 90 95
 Asp Ile Asn Asp Thr Gly Phe Phe Met Asn Leu Glu Glu Asp Met Thr
 100 105 110
 Arg Tyr Ala Tyr Tyr Tyr Ser Gly Ile Gly Ala Gly Val Leu Val Ala
 115 120 125
 Ala Tyr Ile Gln Val Ser Phe Trp Cys Leu Ala Ala Gly Arg Gln Ile
 130 135 140
 His Lys Ile Arg Lys Gln Phe Phe His Ala Ile Met Arg Gln Glu Ile
 145 150 155 160
 Gly Trp Phe Asp Val His Asp Val Gly Glu Leu Asn Thr Arg Leu Thr
 165 170 175
 Asp Asp Val Ser Lys Ile Asn Glu Gly Ile Gly Asp Lys Ile Gly Met
 180 185 190
 Phe Phe Gln Ser Met Ala Thr Phe Phe Thr Gly Phe Ile Val Gly Phe
 195 200 205
 Thr Arg Gly Trp Lys Leu Thr Leu Val Ile Leu Ala Ile Ser Pro Val
 210 215 220
 Leu Gly Leu Ser Ala Ala Val Trp Ala Lys Ile Leu Ser Ser Phe Thr
 225 230 235 240
 Asp Lys Glu Leu Leu Ala Tyr Ala Lys Ala Gly Ala Val Ala Glu Glu
 245 250 255
 Val Leu Ala Ala Ile Arg Thr Val Ile Ala Phe Gly Gly Gln Lys Lys
 260 265 270
 Glu Leu Glu Arg Tyr Asn Lys Asn Leu Glu Glu Ala Lys Arg Ile Gly
 275 280 285
 Ile Lys Lys Ala Ile Thr Ala Asn Ile Ser Ile Gly Ala Ala Phe Leu
 290 295 300
 Leu Ile Tyr Ala Ser Tyr Ala Leu Ala Phe Trp Tyr Gly Thr Thr Leu
 305 310 315 320
 Val Leu Ser Gly Glu Tyr Ser Ile Gly Gln Val Leu Thr Val Phe Phe
 325 330 335

Ser Val Leu Ile Gly Ala Phe Ser Val Gly Gln Ala Ser Pro Ser Ile
 340 345 350
 Glu Ala Phe Ala Asn Ala Arg Gly Ala Ala Tyr Glu Ile Phe Lys Ile
 355 360 365
 Ile Asp Asn Lys Pro Ser Ile Asp Ser Tyr Ser Lys Ser Gly His Lys
 370 375 380
 Pro Asp Asn Ile Lys Gly Asn Leu Glu Phe Arg Asn Val His Phe Ser
 385 390 395 400
 Tyr Pro Ser Arg Lys Glu Val Lys Ile Leu Lys Gly Leu Asn Leu Lys
 405 410 415
 Val Gln Ser Gly Gln Thr Val Ala Leu Val Gly Asn Ser Gly Cys Gly
 420 425 430
 Lys Ser Thr Thr Val Gln Leu Met Gln Arg Leu Tyr Asp Pro Thr Glu
 435 440 445
 Gly Met Val Ser Val Asp Gly Gln Asp Ile Arg Thr Ile Asn Val Arg
 450 455 460
 Phe Leu Arg Glu Ile Ile Gly Val Val Ser Gln Glu Pro Val Leu Phe
 465 470 475 480
 Ala Thr Thr Ile Ala Glu Asn Ile Arg Tyr Gly Arg Glu Asn Val Thr
 485 490 495
 Met Asp Glu Ile Glu Lys Ala Val Lys Glu Ala Asn Ala Tyr Asp Phe
 500 505 510
 Ile Met Lys Leu Pro His Lys Phe Asp Thr Leu Val Gly Glu Arg Gly
 515 520 525
 Ala Gln Leu Ser Gly Gly Gln Lys Gln Arg Ile Ala Ile Ala Arg Ala
 530 535 540
 Leu Val Arg Asn Pro Lys Ile Leu Leu Leu Asp Glu Ala Thr Ser Ala
 545 550 555 560
 Leu Asp Thr Glu Ser Glu Ala Val Val Gln Val Ala Leu Asp Lys Ala
 565 570 575
 Arg Lys Gly Arg Thr Thr Ile Val Ile Ala His Arg Leu Ser Thr Val
 580 585 590
 Arg Asn Ala Asp Val Ile Ala Gly Phe Asp Asp Gly Val Ile Val Glu
 595 600 605
 Lys Gly Asn His Asp Glu Leu Met Lys Glu Lys Gly Ile Tyr Phe Lys
 610 615 620
 Leu Val Thr Met Gln Thr Ala Gly Asn Glu Val Glu Leu Glu Asn Ala
 625 630 635 640
 Ala Asp Glu Ser Lys Ser Glu Ile Asp Ala Leu Glu Met Ser Ser Asn
 645 650 655
 Asp Ser Arg Ser Ser Leu Ile Arg Lys Arg Ser Thr Arg Arg Ser Val
 660 665 670
 Arg Gly Ser Gln Ala Gln Asp Arg Lys Leu Ser Thr Lys Glu Ala Leu
 675 680 685
 Asp Glu Ser Ile Pro Pro Val Ser Phe Trp Arg Ile Met Lys Leu Asn
 690 695 700
 Leu Thr Glu Trp Pro Tyr Phe Val Val Gly Val Phe Cys Ala Ile Ile
 705 710 715 720
 Asn Gly Gly Leu Gln Pro Ala Phe Ala Ile Ile Phe Ser Lys Ile Ile
 725 730 735
 Gly Val Phe Thr Arg Ile Asp Asp Pro Glu Thr Lys Arg Gln Asn Ser
 740 745 750
 Asn Leu Phe Ser Leu Leu Phe Leu Ala Leu Gly Ile Ile Ser Phe Ile
 755 760 765

Thr Phe Phe Leu Gln Gly Phe Thr Phe Gly Lys Ala Gly Glu Ile Leu
 770 775 780
 Thr Lys Arg Leu Arg Tyr Met Val Phe Arg Ser Met Leu Arg Gln Asp
 785 790 795 800
 Val Ser Trp Phe Asp Asp Pro Lys Asn Thr Thr Gly Ala Leu Thr Thr
 805 810 815
 Arg Leu Ala Asn Asp Ala Ala Gln Val Lys Gly Ala Ile Gly Ser Arg
 820 825 830
 Leu Ala Val Ile Thr Gln Asn Ile Ala Asn Leu Gly Thr Gly Ile Ile
 835 840 845
 Ile Ser Phe Ile Tyr Gly Trp Gln Leu Thr Leu Leu Leu Ala Ile
 850 855 860
 Val Pro Ile Ile Ala Ile Ala Gly Val Val Glu Met Lys Met Leu Ser
 865 870 875 880
 Gly Gln Ala Leu Lys Asp Lys Lys Glu Leu Glu Gly Ala Gly Lys Ile
 885 890 895
 Ala Thr Glu Ala Ile Glu Asn Phe Arg Thr Val Val Ser Leu Thr Gln
 900 905 910
 Glu Gln Lys Phe Glu His Met Tyr Ala Gln Ser Leu Gln Val Pro Tyr
 915 920 925
 Arg Asn Ser Leu Arg Lys Ala His Ile Phe Gly Ile Thr Phe Ser Phe
 930 935 940
 Thr Gln Ala Met Met Tyr Phe Ser Tyr Ala Gly Cys Phe Arg Phe Gly
 945 950 955 960
 Ala Tyr Leu Val Ala His Lys Leu Met Ser Phe Glu Asp Val Leu Leu
 965 970 975
 Val Phe Ser Ala Val Val Phe Gly Ala Met Ala Val Gly Gln Val Ser
 980 985 990
 Ser Phe Ala Pro Asp Tyr Ala Lys Ala Lys Ile Ser Ala Ala His Ile
 995 1000 1005
 Ile Met Ile Ile Glu Lys Thr Pro Leu Ile Asp Ser Tyr Ser Thr Glu
 1010 1015 1020
 Gly Leu Met Pro Asn Thr Leu Glu Gly Asn Val Thr Phe Gly Glu Val
 1025 1030 1035 1040
 Val Phe Asn Tyr Pro Thr Arg Pro Asp Ile Pro Val Leu Gln Gly Leu
 1045 1050 1055
 Ser Leu Glu Val Lys Lys Gly Gln Thr Leu Ala Leu Val Gly Ser Ser
 1060 1065 1070
 Gly Cys Gly Lys Ser Thr Val Val Gln Leu Leu Glu Arg Phe Tyr Asp
 1075 1080 1085
 Pro Leu Ala Gly Lys Val Leu Leu Asp Gly Lys Glu Ile Lys Arg Leu
 1090 1095 1100
 Asn Val Gln Trp Leu Arg Ala His Leu Gly Ile Val Ser Gln Glu Pro
 1105 1110 1115 1120
 Ile Leu Phe Asp Cys Ser Ile Ala Glu Asn Ile Ala Tyr Gly Asp Asn
 1125 1130 1135
 Ser Arg Val Val Ser Gln Glu Glu Ile Val Arg Ala Ala Lys Glu Ala
 1140 1145 1150
 Asn Ile His Ala Phe Ile Glu Ser Leu Pro Asn Lys Tyr Ser Thr Lys
 1155 1160 1165
 Val Gly Asp Lys Gly Thr Gln Leu Ser Gly Gly Gln Lys Gln Arg Ile
 1170 1175 1180
 Ala Ile Ala Arg Ala Leu Val Arg Gln Pro His Ile Leu Leu Leu Asp
 1185 1190 1195 1200

Glu Ala Thr Ser Ala Leu Asp Thr Glu Ser Glu Lys Val Val Gln Glu
 1205 1210 1215
 Ala Leu Asp Lys Ala Arg Glu Gly Arg Thr Cys Ile Val Ile Ala His
 1220 1225 1230
 Arg Leu Ser Thr Ile Gln Asn Ala Asp Leu Ile Val Val Phe Gln Asn
 1235 1240 1245
 Gly Arg Val Lys Glu His Gly Thr His Gln Gln Leu Leu Ala Gln Lys
 1250 1255 1260
 Gly Ile Tyr Phe Ser Met Val Ser Val Gln Ala Gly Thr Lys Arg Gln
 1265 1270 1275 1280

<210> 8

<211> 2001

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> 30, 70, 280, 477, 558, 1471, 1651, 1689, 1724

<223> Xaa = Any Amino Acid

<221> VARIANT

<222> 30, 70, 280, 477, 558, 1471, 1651, 1689, 1724

<223> Xaa = Any Amino Acid

<400> 8

Met Ser Ser Leu Gly Phe Thr Ser Lys Glu Gln Arg Asn Leu Gly Leu
 1 5 10 15
 Leu Val His Leu Met Thr Ser Asn Pro Lys Ile Leu Tyr Xaa Pro Ala
 20 25 30
 Gly Ser Glu Val Asp Arg Val Ile Leu Lys Ala Asn Glu Thr Phe Ala
 35 40 45
 Phe Val Gly Asn Val Thr His Tyr Ala Gln Val Trp Leu Asn Ile Ser
 50 55 60
 Ala Glu Ile Arg Ser Xaa Leu Glu Gln Gly Arg Leu Gln Gln His Leu
 65 70 75 80
 Arg Trp Leu Gln Gln Tyr Val Ala Glu Leu Arg Pro His Pro Glu Ala
 85 90 95
 Leu Asn Leu Ser Leu Asp Glu Leu Pro Pro Ala Leu Arg Gln Asp Asn
 100 105 110
 Phe Ser Leu Pro Ser Gly Met Ala Leu Leu Gln Gln Leu Asp Thr Ile
 115 120 125
 Asp Asn Ala Pro Cys Gly Trp Ile Gln Phe Met Ser Lys Val Ser Val
 130 135 140
 Asp Ile Phe Lys Gly Phe Pro Asp Glu Glu Ser Ile Val Asn Tyr Thr
 145 150 155 160
 Leu Asn Gln Ala Tyr Gln Asp Asn Val Thr Val Phe Ala Gly Val Ile
 165 170 175
 Phe Gln Thr Arg Lys Asp Gly Ser Leu Pro Pro His Val His Tyr Lys
 180 185 190
 Ile Arg Gln Asn Ser Ser Phe Thr Glu Lys Thr Asn Glu Ile Arg Arg
 195 200 205
 Ala Tyr Trp Arg Pro Gly Pro Asn Thr Gly Gly Arg Phe Tyr Phe Leu

210 215 220
 Tyr Gly Phe Val Trp Ile Gln Asp Met Met Glu Arg Ala Ile Ile Asp
 225 230 235 240
 Thr Phe Val Gly His Asp Val Val Glu Pro Gly Ser Tyr Val Gln Met
 245 250 255
 Phe Pro Tyr Pro Cys Tyr Thr Arg Asp Asp Phe Leu Phe Val Ile Glu
 260 265 270
 His Met Met Pro Leu Cys Met Xaa Ile Ser Trp Val Tyr Ser Val Ala
 275 280 285
 Met Thr Ile Gln His Ile Val Ala Glu Lys Glu His Arg Leu Lys Glu
 290 295 300
 Val Met Lys Thr Met Gly Leu Asn Asn Ala Val His Trp Val Ala Trp
 305 310 315 320
 Phe Ile Thr Gly Phe Val Gln Leu Ser Ile Ser Val Thr Ala Leu Thr
 325 330 335
 Ala Ile Leu Lys Tyr Gly Gln Val Leu Met His Ser His Val Val Ile
 340 345 350
 Ile Trp Leu Phe Leu Ala Val Tyr Ala Val Ala Thr Ile Met Phe Cys
 355 360 365
 Phe Leu Val Ser Val Leu Tyr Ser Lys Ala Lys Leu Ala Ser Ala Gly
 370 375 380
 Gly Ile Ile Tyr Phe Leu Ser Tyr Val Pro Tyr Met Tyr Val Ala Ile
 385 390 395 400
 Arg Glu Glu Val Ala His Asp Lys Ile Thr Ala Phe Glu Lys Cys Ile
 405 410 415
 Ala Ser Leu Met Ser Thr Thr Ala Phe Gly Leu Gly Ser Lys Tyr Phe
 420 425 430
 Ala Leu Tyr Glu Val Ala Gly Val Gly Ile Gln Trp His Thr Phe Ser
 435 440 445
 Gln Ser Pro Val Glu Gly Asp Asp Phe Asn Leu Leu Leu Ala Val Thr
 450 455 460
 Met Leu Met Val Asp Ala Val Val Tyr Gly Ile Leu Xaa Trp Tyr Ile
 465 470 475 480
 Glu Ala Val His Pro Gly Met Tyr Gly Leu Pro Arg Pro Trp Tyr Phe
 485 490 495
 Pro Leu Gln Lys Ser Tyr Trp Leu Gly Ser Gly Arg Thr Glu Ala Trp
 500 505 510
 Glu Trp Ser Trp Pro Trp Ala Arg Thr Pro Arg Leu Ser Val Met Glu
 515 520 525
 Glu Asp Gln Ala Cys Ala Met Glu Ser Arg Arg Phe Glu Glu Thr Arg
 530 535 540
 Gly Met Glu Glu Glu Pro Thr His Leu Pro Leu Val Val Xaa Val Asp
 545 550 555 560
 Lys Leu Thr Lys Val Tyr Lys Asp Asp Lys Lys Leu Ala Leu Asn Lys
 565 570 575
 Leu Ser Leu Asn Leu Tyr Glu Asn Gln Gly Val Ser Phe Leu Gly His
 580 585 590
 Asn Gly Ala Gly Lys Thr Thr Thr Met Ser Ile Leu Thr Gly Leu Phe
 595 600 605
 Pro Pro Thr Ser Gly Ser Ala Thr Ile Tyr Gly His Asp Ile Arg Thr
 610 615 620
 Glu Met Asp Glu Ile Arg Lys Asn Gly His Val Pro Gln His Asn Val
 625 630 635 640
 Leu Phe Asp Arg Leu Thr Val Glu Glu His Leu Trp Phe Tyr Ser Arg

					645					650				655	
Leu	Lys	Ser	Met	Ala	Gln	Glu	Glu	Ile	Pro	Arg	Glu	Met	Asp	Lys	Met
			660					665					670		
Ile	Glu	Asp	Leu	Glu	Leu	Ser	Asn	Lys	Arg	His	Ser	Leu	Val	Gln	Thr
		675					680					685			
Leu	Ser	Gly	Gly	Met	Lys	Arg	Lys	Val	Ser	Val	Ala	Ile	Ala	Phe	Val
	690					695					700				
Gly	Gly	Ser	Arg	Ala	Ile	Ile	Leu	Asp	Glu	Pro	Thr	Ala	Gly	Val	Asp
705					710					715					720
Pro	Tyr	Ala	Arg	Arg	Ala	Ile	Trp	Asp	Leu	Ile	Leu	Lys	Tyr	Lys	Pro
				725					730					735	
Gly	Arg	Thr	Ile	Leu	Leu	Ser	Thr	His	His	Met	Asp	Glu	Ala	Asp	Leu
			740					745					750		
Leu	Gly	Asp	Arg	Ile	Ala	Ile	Ile	Ser	His	Gly	Lys	Leu	Lys	Cys	Cys
		755					760					765			
Gly	Ser	Pro	Leu	Phe	Leu	Lys	Gly	Thr	Tyr	Gly	Asp	Gly	Tyr	Arg	Leu
	770					775					780				
Thr	Leu	Val	Lys	Arg	Pro	Ala	Glu	Pro	Gly	Gly	Pro	Gln	Glu	Pro	Gly
785					790					795					800
Leu	Ala	Ser	Ser	Pro	Pro	Gly	Arg	Ala	Pro	Leu	Ser	Ser	Cys	Ser	Glu
				805					810					815	
Leu	Gln	Val	Ser	Gln	Phe	Ile	Arg	Lys	His	Val	Ala	Ser	Cys	Leu	Leu
			820					825					830		
Val	Ser	Asp	Thr	Ser	Thr	Glu	Leu	Ser	Tyr	Ile	Leu	Pro	Ser	Glu	Ala
		835					840					845			
Ala	Lys	Lys	Gly	Ala	Phe	Glu	Arg	Leu	Phe	Gln	His	Leu	Glu	Arg	Ser
	850					855					860				
Leu	Asp	Ala	Leu	His	Leu	Ser	Ser	Phe	Gly	Leu	Met	Asp	Thr	Thr	Leu
865					870					875					880
Glu	Glu	Val	Phe	Leu	Lys	Val	Ser	Gly	Gly	Asp	Gln	Ser	Leu	Glu	Asn
				885					890					895	
Ser	Gly	Ala	Asp	Val	Lys	Glu	Ser	Arg	Lys	Asp	Val	Leu	Pro	Gly	Ala
			900					905					910		
Glu	Gly	His	Ala	Ser	Gly	Glu	Gly	His	Ala	Gly	Asn	Leu	Ala	Arg	Cys
		915					920					925			
Ser	Glu	Leu	Thr	Gln	Ser	Gln	Ala	Ser	Leu	Gln	Ser	Ala	Ser	Ser	Val
	930					935					940				
Gly	Ser	Ala	Leu	Gly	Asp	Glu	Gly	Ala	Gly	Tyr	Thr	Asp	Val	Tyr	Gly
945					950					955					960
Asp	Tyr	Pro	Pro	Leu	Phe	Asp	Asn	Pro	Gln	Asp	Pro	Asp	Asn	Val	Ser
				965					970					975	
Leu	Gln	Glu	Val	Glu	Ala	Glu	Ala	Leu	Ser	Arg	Val	Gly	Gln	Gly	Ser
			980					985					990		
Arg	Lys	Leu	Asp	Gly	Gly	Trp	Leu	Lys	Val	Arg	Gln				

1075	1080	1085
Ala Ser Pro Gln Gln Leu Val Ser Thr Phe Arg Leu Pro Ser Gly Val		
1090	1095	1100
Gly Ala Thr Cys Val Leu Lys Ser Pro Ala Asn Gly Ser Leu Gly Pro		
1105	1110	1115
Thr Leu Asn Leu Ser Ser Gly Glu Ser Arg Leu Leu Ala Ala Arg Phe		
1125	1130	1135
Phe Asp Ser Met Cys Leu Glu Ser Phe Thr Gln Gly Leu Pro Leu Ser		
1140	1145	1150
Asn Phe Val Pro Pro Pro Pro Ser Pro Ala Pro Ser Asp Ser Pro Ala		
1155	1160	1165
Ser Pro Asp Glu Asp Leu Gln Ala Trp Asn Val Ser Leu Pro Pro Thr		
1170	1175	1180
Ala Gly Gln Glu Met Trp Thr Ser Ala Pro Ser Leu Pro Arg Leu Val		
1185	1190	1195
Arg Glu Pro Val Arg Cys Thr Cys Ser Ala Gln Gly Thr Gly Phe Ser		
1205	1210	1215
Cys Pro Asn Ser Val Gly Gly His Pro Pro Gln Met Arg Val Val Thr		
1220	1225	1230
Gly Asp Ile Leu Thr Asp Ile Thr Gly His Asn Val Ser Glu Tyr Leu		
1235	1240	1245
Leu Phe Thr Ser Asp Arg Phe Arg Leu His Arg Tyr Gly Ala Ile Thr		
1250	1255	1260
Phe Gly Asn Val Leu Lys Ser Ile Pro Ala Ser Phe Gly Thr Arg Ala		
1265	1270	1275
Pro Pro Met Val Arg Lys Ile Arg Cys Ala Arg Ala Ala Gln Val Phe		
1285	1290	1295
Tyr Asn Asn Lys Gly Tyr His Ser Met Pro Thr Tyr Leu Asn Ser Leu		
1300	1305	1310
Asn Asn Ala Ile Leu Arg Ala Asn Leu Pro Lys Ser Lys Gly Asn Pro		
1315	1320	1325
Ala Ala Tyr Gly Ile Thr Val Thr Asn His Pro Met Asn Lys Thr Ser		
1330	1335	1340
Ala Ser Leu Ser Leu Asp Tyr Leu Leu Gln Gly Thr Asp Val Val Ile		
1345	1350	1355
Ala Ile Phe Ile Ile Val Ala Met Ser Phe Val Pro Ala Ser Phe Val		
1365	1370	1375
Val Phe Leu Val Ala Glu Lys Ser Thr Lys Ala Lys His Leu Gln Phe		
1380	1385	1390
Val Ser Gly Cys Asn Pro Ile Ile Tyr Trp Leu Ala Asn Tyr Val Trp		
1395	1400	1405
Asp Met Leu Asn Tyr Leu Val Pro Ala Thr Cys Cys Val Ile Ile Leu		
1410	1415	1420
Phe Val Phe Asp Leu Pro Ala Tyr Thr Ser Pro Thr Asn Phe Pro Ala		
1425	1430	1435
Val Leu Ser Leu Phe Leu Leu Tyr Gly Trp Ser Ile Thr Pro Ile Met		
1445	1450	1455
Tyr Pro Ala Ser Phe Trp Phe Glu Val Pro Ser Ser Ala Tyr Xaa Phe		
1460	1465	1470
Leu Ile Val Ile Asn Leu Phe Ile Gly Ile Thr Ala Thr Val Ala Thr		
1475	1480	1485
Phe Leu Leu Gln Leu Phe Glu His Asp Lys Asp Leu Lys Val Val Asn		
1490	1495	1500
Ser Tyr Leu Lys Ser Cys Phe Leu Ile Phe Pro Asn Tyr Asn Leu Gly		

1505 1510 1515 1520
 His Gly Leu Met Glu Met Ala Tyr Asn Glu Tyr Ile Asn Glu Tyr Tyr
 1525 1530 1535
 Ala Lys Ile Gly Gln Phe Asp Lys Met Lys Ser Pro Phe Glu Trp Asp
 1540 1545 1550
 Ile Val Thr Arg Gly Leu Val Ala Met Ala Val Glu Gly Val Val Gly
 1555 1560 1565
 Phe Leu Leu Thr Ile Met Cys Gln Tyr Asn Phe Leu Arg Arg Pro Gln
 1570 1575 1580
 Arg Met Pro Val Ser Thr Lys Pro Val Glu Asp Asp Val Asp Val Ala
 1585 1590 1595 1600
 Ser Glu Arg Gln Arg Val Leu Arg Gly Asp Ala Asp Asn Asp Met Val
 1605 1610 1615
 Lys Ile Glu Asn Leu Thr Lys Val Tyr Lys Ser Arg Lys Ile Gly Arg
 1620 1625 1630
 Ile Leu Ala Val Asp Arg Leu Cys Leu Gly Val Arg Pro Gly Glu Cys
 1635 1640 1645
 Phe Gly Xaa Leu Gly Val Asn Gly Ala Gly Lys Thr Ser Thr Phe Lys
 1650 1655 1660
 Met Leu Thr Gly Asp Glu Ser Thr Thr Gly Gly Glu Ala Phe Val Asn
 1665 1670 1675 1680
 Gly His Ser Val Leu Lys Glu Leu Xaa Gln Val Gln Gln Ser Leu Gly
 1685 1690 1695
 Tyr Cys Pro Gln Cys Asp Ala Leu Phe Asp Glu Leu Thr Ala Arg Glu
 1700 1705 1710
 His Leu Gln Leu Tyr Thr Arg Leu Arg Gly Ile Xaa Trp Lys Asp Glu
 1715 1720 1725
 Ala Arg Val Val Lys Trp Ala Leu Glu Lys Leu Glu Leu Thr Lys Tyr
 1730 1735 1740
 Ala Asp Lys Pro Ala Gly Thr Tyr Ser Gly Gly Asn Lys Arg Lys Leu
 1745 1750 1755 1760
 Ser Thr Ala Ile Ala Leu Ile Gly Tyr Pro Ala Phe Ile Phe Leu Asp
 1765 1770 1775
 Glu Pro Thr Thr Gly Met Asp Pro Lys Ala Arg Arg Phe Leu Trp Asn
 1780 1785 1790
 Leu Ile Leu Asp Leu Ile Lys Thr Gly Arg Ser Val Val Leu Thr Ser
 1795 1800 1805
 His Ser Met Glu Glu Cys Glu Ala Leu Cys Thr Arg Leu Ala Ile Met
 1810 1815 1820
 Val Asn Gly Arg Leu Arg Cys Leu Gly Ser Ile Gln His Leu Lys Asn
 1825 1830 1835 1840
 Arg Phe Gly Asp Gly Tyr Met Ile Thr Val Arg Thr Lys Ser Ser Gln
 1845 1850 1855
 Ser Val Lys Asp Val Val Arg Phe Phe Asn Arg Asn Phe Pro Glu Ala
 1860 1865 1870
 Met Leu Lys Glu Arg His His Thr Lys Val Gln Tyr Gln Leu Lys Ser
 1875 1880 1885
 Glu His Ile Ser Leu Ala Gln Val Phe Ser Lys Met Glu Gln Val Ser
 1890 1895 1900
 Gly Val Leu Gly Ile Glu Asp Tyr Ser Val Ser Gln Thr Thr Leu Asp
 1905 1910 1915 1920
 Asn Val Phe Val Asn Phe Ala Lys Lys Gln Ser Asp Asn Leu Glu Gln
 1925 1930 1935
 Gln Glu Thr Glu Pro Pro Ser Ala Leu Gln Ser Pro Leu Gly Cys Leu

1940	1945	1950
Leu Ser Leu Leu Arg Pro Arg Ser Ala Pro Thr Glu Leu Arg Ala Leu		
1955	1960	1965
Val Ala Asp Glu Pro Glu Asp Leu Asp Thr Glu Asp Glu Gly Leu Ile		
1970	1975	1980
Ser Phe Glu Glu Glu Arg Ala Gln Leu Ser Phe Asn Thr Asp Thr Leu		
1985	1990	1995
Cys		2000

<210> 9
 <211> 646
 <212> PRT
 <213> Homo sapiens

<400> 9

Met	Ala	Glu	Lys	Ala	Leu	Glu	Ala	Val	Gly	Cys	Gly	Leu	Gly	Pro	Gly
1			5						10					15	
Ala	Val	Ala	Met	Ala	Val	Thr	Leu	Glu	Asp	Gly	Ala	Glu	Pro	Pro	Val
		20						25					30		
Leu	Thr	Thr	His	Leu	Lys	Lys	Val	Glu	Asn	His	Ile	Thr	Glu	Ala	Gln
		35					40					45			
Arg	Phe	Ser	His	Leu	Pro	Lys	Arg	Ser	Ala	Val	Asp	Ile	Glu	Phe	Val
	50					55					60				
Glu	Leu	Ser	Tyr	Ser	Val	Arg	Glu	Gly	Pro	Cys	Trp	Arg	Lys	Arg	Gly
65					70					75					80
Tyr	Lys	Thr	Leu	Leu	Lys	Cys	Leu	Ser	Gly	Lys	Phe	Cys	Arg	Arg	Glu
			85						90					95	
Leu	Ile	Gly	Ile	Met	Gly	Pro	Ser	Gly	Ala	Gly	Lys	Ser	Thr	Phe	Met
			100					105					110		
Asn	Ile	Leu	Ala	Gly	Tyr	Arg	Glu	Ser	Gly	Met	Lys	Gly	Gln	Ile	Leu
		115					120					125			
Val	Asn	Gly	Arg	Pro	Arg	Glu	Leu	Arg	Thr	Phe	Arg	Lys	Met	Ser	Cys
	130					135					140				
Tyr	Ile	Met	Gln	Asp	Asp	Met	Leu	Leu	Pro	His	Leu	Thr	Val	Leu	Glu
145					150					155					160
Ala	Met	Met	Val	Ser	Ala	Asn	Leu	Asn	Leu	Thr	Glu	Asn	Pro	Asp	Val
			165						170					175	
Lys	Asn	Asp	Leu	Val	Thr	Glu	Ile	Leu	Thr	Ala	Leu	Gly	Leu	Met	Ser
		180						185					190		
Cys	Ser	His	Thr	Arg	Thr	Ala	Leu	Leu	Ser	Gly	Gly	Gln	Arg	Lys	Arg
		195					200					205			
Leu	Ala	Ile	Ala	Leu	Glu	Leu	Val	Asn	Asn	Pro	Pro	Val	Met	Phe	Phe
	210					215					220				
Asp	Glu	Pro	Thr	Ser	Gly	Leu	Asp	Ser	Ala	Ser	Cys	Phe	Gln	Val	Val
225					230					235					240
Ser	Leu	Met	Lys	Ser	Leu	Ala	Gln	Gly	Gly	Arg	Thr	Ile	Ile	Cys	Thr
			245						250					255	
Ile	His	Gln	Pro	Ser	Ala	Lys	Leu	Phe	Glu	Met	Phe	Asp	Lys	Leu	Tyr
		260					265					270			
Ile	Leu	Ser	Gln	Gly	Gln	Cys	Ile	Phe	Lys	Gly	Val	Val	Thr	Asn	Leu
	275						280					285			
Ile	Pro	Tyr	Leu	Lys	Gly	Leu	Gly	Leu	His	Cys	Pro	Thr	Tyr	His	Asn

290 295 300
 Pro Ala Asp Phe Ile Ile Glu Val Ala Ser Gly Glu Tyr Gly Asp Leu
 305 310 315 320
 Asn Pro Met Leu Phe Arg Ala Val Gln Asn Gly Leu Cys Ala Met Ala
 325 330 335
 Glu Lys Lys Ser Ser Pro Glu Lys Asn Glu Val Pro Ala Pro Cys Pro
 340 345 350
 Pro Cys Pro Pro Glu Val Asp Pro Ile Glu Ser His Thr Phe Ala Thr
 355 360 365
 Ser Thr Leu Thr Gln Phe Cys Ile Leu Phe Lys Arg Thr Phe Leu Ser
 370 375 380
 Ile Leu Arg Asp Thr Val Leu Thr His Leu Arg Phe Met Ser His Val
 385 390 395 400
 Val Ile Gly Val Leu Ile Gly Leu Leu Tyr Leu His Ile Gly Asp Asp
 405 410 415
 Ala Ser Lys Val Phe Asn Asn Thr Gly Cys Leu Phe Phe Ser Met Leu
 420 425 430
 Phe Leu Met Phe Ala Ala Leu Met Pro Thr Val Leu Thr Phe Pro Leu
 435 440 445
 Glu Met Ala Val Phe Met Arg Glu His Leu Asn Tyr Trp Tyr Ser Leu
 450 455 460
 Lys Ala Tyr Tyr Leu Ala Lys Thr Met Ala Asp Val Pro Phe Gln Val
 465 470 475 480
 Val Cys Pro Val Val Tyr Cys Ser Ile Val Tyr Trp Met Thr Gly Gln
 485 490 495
 Pro Ala Glu Thr Ser Arg Phe Leu Leu Phe Ser Ala Leu Ala Thr Ala
 500 505 510
 Thr Ala Leu Val Ala Gln Ser Leu Gly Leu Leu Ile Gly Ala Ala Ser
 515 520 525
 Asn Ser Leu Gln Val Ala Thr Phe Val Gly Pro Val Thr Ala Ile Pro
 530 535 540
 Val Leu Leu Phe Ser Gly Phe Phe Val Ser Phe Lys Thr Ile Pro Thr
 545 550 555 560
 Tyr Leu Gln Trp Ser Ser Tyr Leu Ser Tyr Val Arg Tyr Gly Phe Glu
 565 570 575
 Gly Val Ile Leu Thr Ile Tyr Gly Met Glu Arg Gly Asp Leu Thr Cys
 580 585 590
 Leu Glu Glu Arg Cys Pro Phe Arg Glu Pro Gln Ser Ile Leu Arg Ala
 595 600 605
 Leu Asp Val Glu Asp Ala Lys Leu Tyr Met Asp Phe Leu Val Leu Gly
 610 615 620
 Ile Phe Phe Leu Ala Leu Arg Leu Leu Ala Tyr Leu Val Leu Arg Tyr
 625 630 635 640
 Arg Val Lys Ser Glu Arg
 645

<210> 10

<211> 638

<212> PRT

<213> Homo sapiens

<400> 10

Met Glu Ala Thr Glu Thr Asp Leu Leu Asn Gly His Leu Lys Lys Val

1	5	10	15
Asp Asn Asn Leu Thr Glu Ala Gln Arg Phe Ser Ser Leu Pro Arg Arg			
20	25	30	
Ala Ala Val Asn Ile Glu Phe Arg Asp Leu Ser Tyr Ser Val Pro Glu			
35	40	45	
Gly Pro Trp Trp Arg Lys Lys Gly Tyr Lys Thr Leu Leu Lys Gly Ile			
50	55	60	
Ser Gly Lys Phe Asn Ser Gly Glu Leu Val Ala Ile Met Gly Pro Ser			
65	70	75	80
Gly Ala Gly Lys Ser Thr Leu Met Asn Ile Leu Ala Gly Tyr Arg Glu			
85	90	95	
Thr Gly Met Lys Gly Ala Val Leu Ile Asn Gly Leu Pro Arg Asp Leu			
100	105	110	
Arg Cys Phe Arg Lys Val Ser Cys Tyr Ile Met Gln Asp Asp Met Leu			
115	120	125	
Leu Pro His Leu Thr Val Gln Glu Ala Met Met Val Ser Ala His Leu			
130	135	140	
Lys Leu Gln Glu Lys Asp Glu Gly Arg Arg Glu Met Val Lys Glu Ile			
145	150	155	160
Leu Thr Ala Leu Gly Leu Leu Ser Cys Ala Asn Thr Arg Thr Gly Ser			
165	170	175	
Leu Ser Gly Gly Gln Arg Lys Arg Leu Ala Ile Ala Leu Glu Leu Val			
180	185	190	
Asn Asn Pro Pro Val Met Phe Phe Asp Glu Pro Thr Ser Gly Leu Asp			
195	200	205	
Ser Ala Ser Cys Phe Gln Val Val Ser Leu Met Lys Gly Leu Ala Gln			
210	215	220	
Gly Gly Arg Ser Ile Ile Cys Thr Ile His Gln Pro Ser Ala Lys Leu			
225	230	235	240
Phe Glu Leu Phe Asp Gln Leu Tyr Val Leu Ser Gln Gly Gln Cys Val			
245	250	255	
Tyr Arg Gly Lys Val Cys Asn Leu Val Pro Tyr Leu Arg Asp Leu Gly			
260	265	270	
Leu Asn Cys Pro Thr Tyr His Asn Pro Ala Asp Phe Val Met Glu Val			
275	280	285	
Ala Ser Gly Glu Tyr Gly Asp Gln Asn Ser Arg Leu Val Arg Ala Val			
290	295	300	
Arg Glu Gly Met Cys Asp Ser Asp His Lys Arg Asp Leu Gly Gly Asp			
305	310	315	320
Ala Glu Val Asn Pro Phe Leu Trp His Arg Pro Ser Glu Glu Val Lys			
325	330	335	
Gln Thr Lys Arg Leu Lys Gly Leu Arg Lys Asp Ser Ser Ser Met Glu			
340	345	350	
Gly Cys His Ser Phe Ser Ala Ser Cys Leu Thr Gln Phe Cys Ile Leu			
355	360	365	
Phe Lys Arg Thr Phe Leu Ser Ile Met Arg Asp Ser Val Leu Thr His			
370	375	380	
Leu Arg Ile Thr Ser His Ile Gly Ile Gly Leu Leu Ile Gly Leu Leu			
385	390	395	400
Tyr Leu Gly Ile Gly Asn Glu Ala Lys Lys Val Leu Ser Asn Ser Gly			
405	410	415	
Phe Leu Phe Phe Ser Met Leu Phe Leu Met Phe Ala Ala Leu Met Pro			
420	425	430	
Thr Val Leu Thr Phe Pro Leu Glu Met Gly Val Phe Leu Arg Glu His			

435	440	445
Leu Asn Tyr Trp Tyr Ser Leu Lys Ala Tyr Tyr Leu Ala Lys Thr Met		
450	455	460
Ala Asp Val Pro Phe Gln Ile Met Phe Pro Val Ala Tyr Cys Ser Ile		
465	470	475
Val Tyr Trp Met Thr Ser Gln Pro Ser Asp Ala Val Ala Phe Val Leu		480
	485	490
Phe Ala Ala Leu Gly Thr Met Thr Ser Leu Val Ala Gln Ser Leu Gly		495
	500	505
Leu Leu Ile Gly Ala Ala Ser Thr Ser Leu Gln Val Ala Thr Phe Val		510
	515	520
Gly Pro Val Thr Ala Ile Pro Val Leu Leu Phe Ser Gly Phe Phe Val		525
	530	535
Ser Phe Asp Thr Ile Pro Thr Tyr Leu Gln Trp Met Ser Tyr Ile Ser		540
545	550	555
Tyr Val Arg Tyr Gly Phe Glu Gly Val Ile Leu Ser Ile Tyr Gly Leu		560
	565	570
Asp Arg Glu Asp Leu His Cys Asp Ile Asp Glu Thr Cys His Phe Gln		575
	580	585
Lys Ser Glu Ala Ile Leu Arg Glu Leu Asp Val Glu Asn Ala Lys Leu		590
	595	600
Tyr Leu Asp Phe Ile Val Leu Gly Ile Phe Phe Ile Ser Leu Arg Leu		605
	610	615
Ile Ala Tyr Phe Val Leu Arg Tyr Lys Ile Arg Ala Glu Arg		620
625	630	635